## STIG Biotechnology Systems Branch

## RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:

Source:

Date Processed by STIC:

10 533, 158

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,

2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE <u>CHECKER VERSION 4.2.2 PROGRAM</u>, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- 3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 01/14/05):
  U.S. Patent and Trademark Office, Mail Stop Sequence, Customer Window, Randolph Building, 401 Dulany Street, Alexandria, VA 22314

Revised 01/24/05



PCT

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/533,158

DATE: 05/12/2005

TIME: 08:47:25

Input Set : A:\Sequence Listings - Atty Dckt 7388-84325.txt

Output Set: N:\CRF4\05122005\J533158.raw

4 <110> APPLICANT: Hisamitsu Pharmaceutical Co., Inc.

Chiba-Prefecture

7 <120> TITLE OF INVENTION: Nucleic acids isolated from stage 4s neuroblastoma

9 <130> FILE REFERENCE: FP03-0298-00

C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/533,158

C--> 11 <141> CURRENT FILING DATE: 2005-04-28

11 <150> PRIOR APPLICATION NUMBER: JP 2002-316586

12 <151> PRIOR FILING DATE: 2002-10-30

14 <160> NUMBER OF SEQ ID NOS: 1082

16 <170> SOFTWARE: PatentIn Ver. 2.1

Does Not Comply

Corrected Diskette Needed

(PS.1)

## ERRORED SEQUENCES

18684 <210> SEQ ID NO: 1074

18685 <211> LENGTH 20

18686 <212> TYPE: DNA

18687 <213> ORGANISM: Artificial Sequence

18689 <220> FEATURE:

18690 <223> OTHER INFORMATION: Synthetic Primer: nbla23300-lr

18692 <400> SEQUENCE: 1074

pls group nucleotides in sets of 10s E--> 18693 acatcaac ttccctccaa 18696 <del>≥210> SEQ ID NQ: 107</del>5

18697 <211> LENGTH: (21

18698 <212> TYPE: DNA

18699 <213> ORGANISM: Artificial Sequence

18701 <220> FEATURE:

18702 <223> OTHER INFORMATION: Synthetic Primer: nbla23664-1f

18704 <400> SEQUENCE: 1075

E--> 18705 ctttcatttc tcctgctgtc c

For exist ecceptatua 20 acatyst ccar etcossatua 20

5/12/05

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/533,158

DATE: 05/12/2005 TIME: 08:47:28

.

Input Set : A:\Sequence Listings - Atty Dckt 7388-84325.txt

Output Set: N:\CRF4\05122005\J533158.raw

L:11 M:270 C: Current Application Number differs, Replaced Current Application No L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date L:8675 M:283 W: Missing Blank Line separator, <400> field identifier L:11302 M:283 W: Missing Blank Line separator, <400> field identifier L:11589 M:283 W: Missing Blank Line separator, <400> field identifier L:12008 M:283 W: Missing Blank Line separator, <400> field identifier

L:18693 M:254 E: No. of Bases conflict, LENGTH:Input:20 Counted:18 SEQ:1074 L:18693 M:252 E: No. of Seq. differs, <211> LENGTH:Input:20 Found:18 SEQ:1074

L:18705 M:254 E: No. of Bases conflict, LENGTH:Input:20 Counted:21 SEQ:1075